

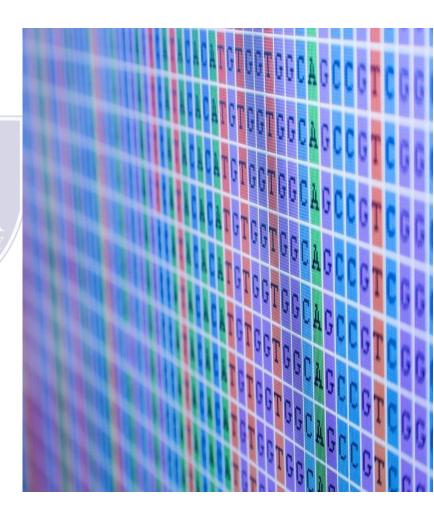
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FACULTY OF ENGINEERING & TECHNOLOGY DEPARTMENT OF BIOTECHNOLOGY

• What is DNASequencing?

History of development A

 Basic Methods- Chain termination and Chemical modification



method

• Determining the precise order of nucleotides in DNA.



• We need to determine the order of nucleotide bases in a strand of DNA for sequencing.

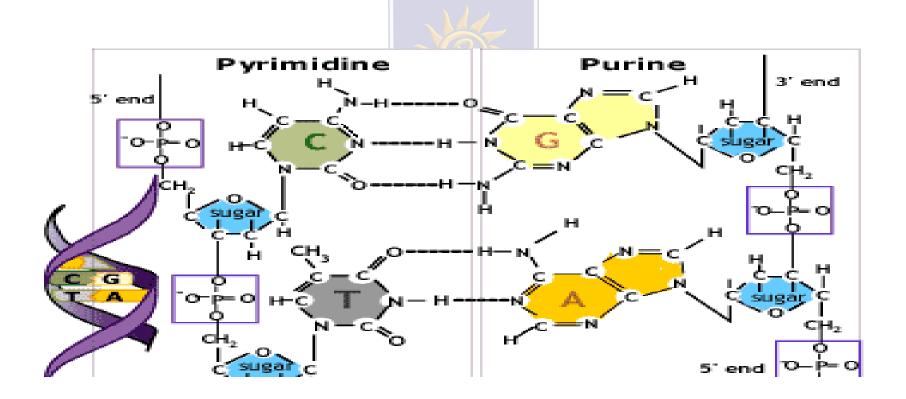
The Need for DNA Sequencing

- Geneisolation
- Sequence charaterization
- Forensics
- MolecularArcheology
- Gene Gene Interaction
- Gene Protein Interaction
- Cloning



DNA

- Deoxyribonucleic Acid Stores genetic information
 Four different nucleotides A,T,G,C
- DNA comprises of a long molecule analogous to a chain, while the links of the chain are called Nucleotides



- 1870 Miescher discovers DNA
- 1940 Avery: Proposes DNA as 'Genetic Material' 1953 -

Watson & Crick "double helical structure" 1970 - Wu:

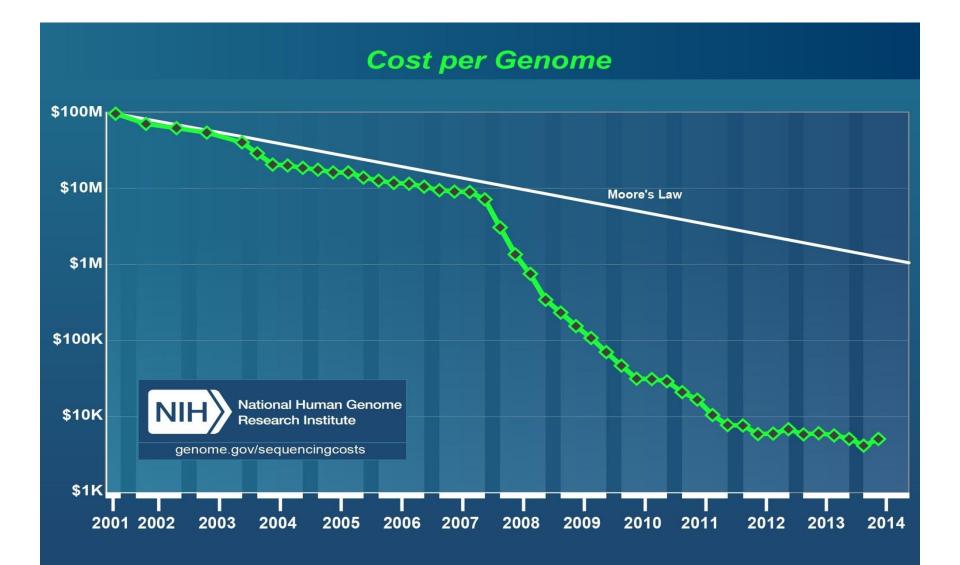
Sequences λ Cohesive End DNA **1977** – Sanger: Dideoxy

Chain Termination

- 1977 Gilbert: Chemical Degradation
- 1986 Partial Automation
- 1990 Cycle Sequencing, Improved Sequencing Enzymes,

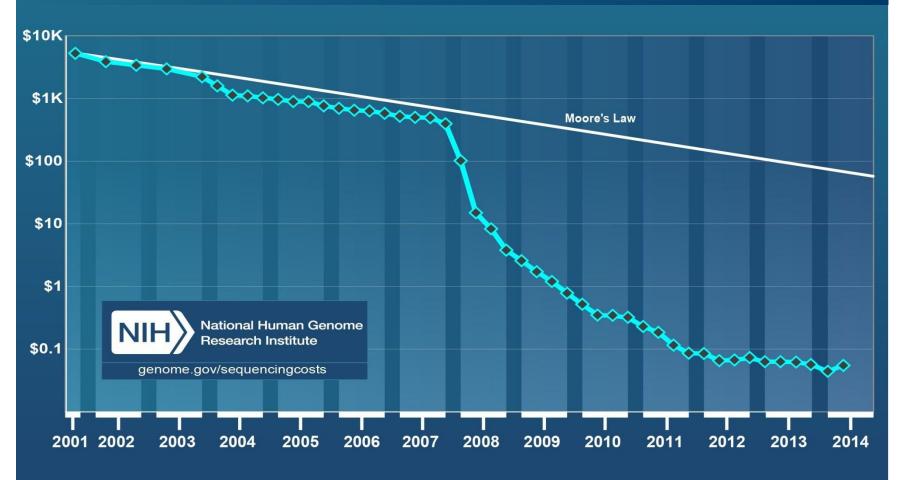
Improved fluorescent detection schemes

2002 - NGS: 454, pyro sequencing



Cost per Megabases

Cost per Raw Megabase of DNA Sequence



- To determine the order of the nucleotide bases adenine, guanine, cytosine, and thymine in a molecule of DNA two methods were used
 - 1. Maxam and Gilbert; Chemical Sequencing
 - 2. Sanger; Chain Termination Sequencing
- These two are conventional methods
- Robotics and automated sequencing are based on these methods

- In 1976–1977, Allan Maxam and Walter Gilbert developed a DNA sequencing method based on chemical modification of DNA and subsequent cleavage at specific bases
 - I. Chemical Modification of DNA; radioactive labeling at one 5' end of the DNA (typically by a kinase reaction using gamma-³²P ATP)
 - II. Purification of the DNA fragment to be sequenced
 - III. Chemical treatment generates breaks in DNA
 - IV. Run on the gel